



## SEQUENCE LISTING

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Cheng, Xing  
Zhou, Helen

<120> Functional Mutations in Respiratory Syncytial Virus

<130> 7682-132-999

<140> US 10/672,302  
<141> 2003-09-26

<160> 92

<170> PatentIn version 3.1

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50 55 60  
Glu Leu Asp Arg Thr Glu Glu Tyr Ala Leu Gly Val Val Gly Val Leu  
65 70 75 80  
Glu Ser Tyr Ile Gly Ser Ile Asn Asn Ile Thr Lys Gln Ser Ala Cys  
85 90 95  
Val Ala Met Ser Lys Leu Leu Thr Glu Leu Asn Ser Asp Asp Ile Lys  
100 105 110  
Lys Leu Arg Asp Asn Glu Glu Leu Asn Ser Pro Lys Ile Arg Val Tyr  
115 120 125  
Asn Thr Val Ile Ser Tyr Ile Glu Ser Asn Arg Lys Asn Asn Lys Gln  
130 135 140  
Thr Ile His Leu Leu Lys Arg Leu Pro Ala Asp Val Leu Lys Lys Thr  
145 150 155 160  
Ile Lys Asn Thr Leu Asp Ile His Lys Ser Ile Thr Ile Asn Asn Pro  
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50 55 60  
Pro Gln Arg Thr Ala Glu Tyr Ala Leu Gly Thr Ile Gly Val Leu Lys  
65 70 75 80  
Ser Tyr Leu Glu Lys Thr Asn Asn Ile Thr Lys Ser Ile Ala Cys Gly  
85 90 95  
Ser Leu Ile Thr Val Leu Gln Asn Leu Asp Val Gly Leu Val Ile Gln  
100 105 110  
Ala Arg Asp Ser Asn Thr Glu Asp Thr Asn Tyr Leu Arg Ser Cys Asn  
115 120 125  
Thr Ile Leu Ser Tyr Ile Asp Lys Ile His Lys Lys Arg Gln Ile Ile  
130 135 140  
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165 170 175

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Gly Arg Asn Cys Lys Tyr Ser His Asn Tyr Trp Glu Trp  
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1 5 10 15  
Gly Arg Asn Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
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Gly Arg Asn Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
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Gly Arg Asn Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
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1 5 10 15

Gly Arg Arg Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
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<400> 35  
Met Ser Arg Arg Pro Cys Lys Phe Glu Val Gln Gly Phe Cys Leu Asn  
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Gly Arg Asn Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
20 25

<210> 36  
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<400> 36  
Met Ser Val Arg Pro Cys Lys Phe Glu Val Arg Gly Phe Cys Leu Asn  
1 5 10 15

Gly Arg Asn Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
20 25

<210> 37  
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<400> 38  
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1 5 10 15  
Gly Arg Asn Cys Lys Tyr Ser His Lys Tyr Trp Glu Trp  
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1 5 10 15  
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Arg	Gly	Lys	Arg	Cys	His	Phe	Ser	His	Asn	Tyr	Phe	Glu	Trp		
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Arg	Gly	Lys	Arg	Cys	His	Phe	Ser	His	Asn	Tyr	Phe	Glu	Trp		
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<400> 43															
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Asn	Gly	Lys	Arg	Cys	His	Phe	Ser	His	Asn	Tyr	Phe	Glu	Trp		
								20		25			30		

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<400> 44															
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Asn Gly Lys Arg Cys His Phe Ser His Asn Tyr Phe Glu Trp  
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Asn Gly Lys Arg Cys His Phe Ser His Asn Tyr Phe Glu Trp  
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Met Ser Arg Arg Asn Pro Cys Lys Phe Glu Ile Gln Gly His Cys Leu  
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Asn Gly Lys Asn Cys His Phe Ser His Asn Tyr Phe Glu Trp  
20 25 30

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<400> 49  
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1 5 10 15  
Asn Gly Lys Asn Cys His Phe Ser His Asn Tyr Phe Glu Trp  
20 25 30

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<210> 52  
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<223> mutant of human RSV A2 P, around residue 176

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gaagatatg

9

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<213> Human respiratory syncytial virus strain A2

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<210> 55  
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<210> 56  
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<213> Human respiratory syncytial virus strain long

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<210> 57  
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<210> 58  
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<210> 59  
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<210> 61  
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1 5 10

<210> 62  
<211> 16  
<212> PRT  
<213> Bovine respiratory syncytial virus

<400> 62  
Ile Glu Thr Phe Asp Asn Asn Glu Glu Glu Ser Ser Tyr Ser Asp Glu  
1 5 10 15

<210> 63  
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<213> Bovine respiratory syncytial virus

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<210> 64  
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1 5 10 15

<210> 65  
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<400> 65

Ile Val Glu Asp Glu Ser Thr Ser Gly Glu Ser Glu Glu  
1 5 10

<210> 66  
<211> 16  
<212> PRT  
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Ile Glu Thr Phe Asp Asn Asn Glu Glu Glu Ser Ser Tyr Ser Asp Glu  
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<210> 67  
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<210> 68  
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<212> PRT  
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<220>  
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<400> 68  
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<210> 69  
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<220>  
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<400> 69  
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<210> 70  
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<220>  
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<400> 70  
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<210> 71  
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<220>  
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<400> 71  
Leu Leu Glu Gly Asn Asp Ser Asp Asn Asp Leu Ser Leu Glu Phe  
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<210> 72  
<211> 16  
<212> PRT  
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<220>  
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<400> 72  
Ile Glu Thr Phe Asp Asn Asn Glu Glu Glu Ser Ser Tyr Ser Glu Glu  
1 5 10 15

<210> 73  
<211> 15  
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<220>  
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<400> 73  
Leu Leu Glu Gly Asn Asp Asp Asp Asn Asp Leu Asp Leu Glu Phe  
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<210> 74  
<211> 16  
<212> PRT  
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<220>  
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<400> 74  
Ile Glu Thr Phe Asp Asn Asn Glu Glu Glu Ser Ser Tyr Ser Glu Glu  
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<210> 75  
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<210> 76  
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<220>  
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<400> 76  
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<210> 77  
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<212> PRT  
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<220>  
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<400> 77  
Leu Leu Glu Gly Asn Asp Ala Asp Asn Asp Leu Ala Leu Glu Phe  
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<210> 78  
<211> 16  
<212> PRT  
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<220>  
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<400> 78  
Ile Glu Thr Phe Asp Asn Asn Glu Glu Glu Leu Arg Tyr Leu Glu Glu  
1 5 10 15

<210> 79  
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<400> 79  
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<213> Human respiratory syncytial virus strain A2  
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<210> 81  
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<220>  
 <223> example conservatively substituted variation of RSV A2 P residues  
 171-176

<400> 81  
 Val Gly Ile Lys Asp Asp  
 1 5

<210> 82  
 <211> 6  
 <212> PRT  
 <213> artificial

<220>  
 <223> example conservatively substituted variation of RSV A2 P residues  
 171-176

<400> 82  
 Ile Gly Val Lys Asp Glu  
 1 5

<210> 83  
 <211> 241  
 <212> PRT  
 <213> Human respiratory syncytial virus strain A2

<400> 83  
 Met Glu Lys Phe Ala Pro Glu Phe His Gly Glu Asp Ala Asn Asn Arg  
 1 5 10 15  
 Ala Thr Lys Phe Leu Glu Ser Ile Lys Gly Lys Phe Thr Ser Pro Lys  
 20 25 30  
 Asp Pro Lys Lys Asp Ser Ile Ile Ser Val Asn Ser Ile Asp Ile  
 35 40 45  
 Glu Val Thr Lys Glu Ser Pro Ile Thr Ser Asn Ser Thr Ile Ile Asn  
 50 55 60  
 Pro Thr Asn Glu Thr Asp Asp Thr Ala Gly Asn Lys Pro Asn Tyr Gln  
 65 70 75 80  
 Arg Lys Pro Leu Val Ser Phe Lys Glu Asp Pro Thr Pro Ser Asp Asn  
 85 90 95  
 Pro Phe Ser Lys Leu Tyr Lys Glu Thr Ile Glu Thr Phe Asp Asn Asn  
 100 105 110  
 Glu Glu Glu Ser Ser Tyr Ser Tyr Glu Glu Ile Asn Asp Gln Thr Asn  
 115 120 125  
 Asp Asn Ile Thr Ala Arg Leu Asp Arg Ile Asp Glu Lys Leu Ser Glu  
 130 135 140  
 Ile Leu Gly Met Leu His Thr Leu Val Val Ala Ser Ala Gly Pro Thr  
 145 150 155 160

Ser Ala Arg Asp Gly Ile Arg Asp Ala Met Ile Gly Leu Arg Glu Glu  
165 170 175  
Met Ile Glu Lys Ile Arg Thr Glu Ala Leu Met Thr Asn Asp Arg Leu  
180 185 190  
Glu Ala Met Ala Arg Leu Arg Asn Glu Ser Glu Lys Met Ala Lys  
195 200 205  
Asp Thr Ser Asp Glu Val Ser Leu Asn Pro Thr Ser Glu Lys Leu Asn  
210 215 220  
Asn Leu Leu Glu Gly Asn Asp Ser Asp Asn Asp Leu Ser Leu Glu Asp  
225 230 235 240  
Phe

<210> 84  
<211> 90  
<212> PRT  
<213> Human respiratory syncytial virus strain A2

<400> 84  
Met Thr Met Pro Lys Ile Met Ile Leu Pro Asp Lys Tyr Pro Cys Ser  
1 5 10 15  
Ile Thr Ser Ile Leu Ile Thr Ser Arg Cys Arg Val Thr Met Tyr Asn  
20 25 30  
Gln Lys Asn Thr Leu Cys Leu Asn Gln Asn Asn Pro Asn Asn His Met  
35 40 45  
Tyr Ser Pro Asn Gln Thr Phe Asn Glu Ile His Trp Thr Ser Gln Glu  
50 55 60  
Leu Ile Asp Thr Ile Gln Asn Phe Leu Gln His Leu Gly Ile Ile Glu  
65 70 75 80  
Asp Ile Tyr Thr Ile Tyr Ile Leu Val Ser  
85 90

<210> 85  
<211> 5  
<212> PRT  
<213> P protein phosphorylation mutant 1

<400> 85  
Leu Arg Leu Ser Ser  
1 5

<210> 86  
<211> 5  
<212> PRT  
<213> P protein phosphorylation mutant 2

<400> 86  
Asp Asp Asp Ser Ser  
1 5

<210> 87  
<211> 5  
<212> PRT  
<213> P protein phosphorylation mutant 3

<400> 87  
Ser Ser Ser Asp Asp  
1 5

<210> 88

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<211> 5
<212> PRT
<213> P protein phosphorylation mutant 4

<400> 88
Ser Ser Ser Ala Ala
1 5

<210> 89
<211> 5
<212> PRT
<213> P protein phosphorylation mutant 5

<400> 89
Leu Arg Leu Ala Ala
1 5

<210> 90
<211> 5
<212> PRT
<213> P protein phosphorylation mutant 6

<400> 90
Leu Arg Leu Asp Asp
1 5

<210> 91
<211> 6
<212> DNA
<213> artificial

<220>
<223> KpnI restriction site

<400> 91
ggtacc 6

<210> 92
<211> 32
<212> DNA
<213> artificial

<220>
<223> synthetic oligonucleotide

<400> 92
agttacttaa aaagaggggc aaataaggta cc 32

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